

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/580,076
Source: IFWP
Date Processed by STIC: 05/31/2006

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 05/31/2006

PATENT APPLICATION: US/10/580,076

TIME: 13:56:01

Input Set: A:\8764.000062.txt

Output Set: N:\CRF4\05312006\J580076.raw

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3 <110> APPLICANT: Bayer BioScience N.V.
4     D'Halluin, Kathleen
5     Vanderstraeten, Chantal
6     Ruiter, Rene
8 <120> TITLE OF INVENTION: Improved targeted DNA insertion in plants
10 <130> FILE REFERENCE: BCS 03-2007 WO1
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/580,076
C--> 12 <141> CURRENT FILING DATE: 2006-05-18
12 <150> PRIOR APPLICATION NUMBER: EP 03078700.6
13 <151> PRIOR FILING DATE: 2003-11-18
15 <160> NUMBER OF SEQ ID NOS: 7
17 <170> SOFTWARE: PatentIn version 3.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 244
21 <212> TYPE: PRT
22 <213> ORGANISM: Saccharomyces cerevisiae
24 <400> SEQUENCE: 1
26 Met Ala Lys Pro Pro Lys Lys Lys Arg Lys Val Asn Ile Lys Lys Asn
27 1          5          10          15
29 Gln Val Met Asn Leu Gly Pro Asn Ser Lys Leu Leu Lys Glu Tyr Lys
30          20          25          30
32 Ser Gln Leu Ile Glu Leu Asn Ile Glu Gln Phe Glu Ala Gly Ile Gly
33          35          40          45
35 Leu Ile Leu Gly Asp Ala Tyr Ile Arg Ser Arg Asp Glu Gly Lys Thr
36          50          55          60
38 Tyr Cys Met Gln Phe Glu Trp Lys Asn Lys Ala Tyr Met Asp His Val
39 65          70          75          80
41 Cys Leu Leu Tyr Asp Gln Trp Val Leu Ser Pro Pro His Lys Lys Glu
42          85          90          95
44 Arg Val Asn His Leu Gly Asn Leu Val Ile Thr Trp Gly Ala Gln Thr
45          100         105         110
47 Phe Lys His Gln Ala Phe Asn Lys Leu Ala Asn Leu Phe Ile Val Asn
48          115         120         125
50 Asn Lys Lys Thr Ile Pro Asn Asn Leu Val Glu Asn Tyr Leu Thr Pro
51          130         135         140
53 Met Ser Leu Ala Tyr Trp Phe Met Asp Asp Gly Gly Lys Trp Asp Tyr
54 145         150         155         160
56 Asn Lys Asn Ser Thr Asn Lys Ser Ile Val Leu Asn Thr Gln Ser Phe
57          165         170         175
59 Thr Phe Glu Glu Val Glu Tyr Leu Val Lys Gly Leu Arg Asn Lys Phe
60          180         185         190
62 Gln Leu Asn Cys Tyr Val Lys Ile Asn Lys Asn Lys Pro Ile Ile Tyr
63          195         200         205

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```

65 Ile Asp Ser Met Ser Tyr Leu Ile Phe Tyr Asn Leu Ile Lys Pro Tyr
66      210                      215                      220
68 Leu Ile Pro Gln Met Met Tyr Lys Leu Pro Asn Thr Ile Ser Ser Glu
69 225                      230                      235                      240
71 Thr Phe Leu Lys
74 <210> SEQ ID NO: 2
75 <211> LENGTH: 732
76 <212> TYPE: DNA
77 <213> ORGANISM: Artificial
79 <220> FEATURE:
80 <223> OTHER INFORMATION: synthetic DNA sequence encoding I-SceI (UIPAC code)
82 <220> FEATURE:
83 <221> NAME/KEY: misc_feature
84 <222> LOCATION: (6)..(6)
85 <223> OTHER INFORMATION: N=A,G,C or T
88 <220> FEATURE:
89 <221> NAME/KEY: variation
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91 <223> OTHER INFORMATION: AGR
94 <220> FEATURE:
95 <221> NAME/KEY: variation
96 <222> LOCATION: (61)..(63)
97 <223> OTHER INFORMATION: TTR
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102 <222> LOCATION: (73)..(75)
103 <223> OTHER INFORMATION: AGY
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108 <222> LOCATION: (79)..(81)
109 <223> OTHER INFORMATION: TTR
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114 <222> LOCATION: (82)..(84)
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118 <220> FEATURE:
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120 <222> LOCATION: (97)..(99)
121 <223> OTHER INFORMATION: AGY
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126 <222> LOCATION: (103)..(105)
127 <223> OTHER INFORMATION: TTR
130 <220> FEATURE:
131 <221> NAME/KEY: variation
132 <222> LOCATION: (112)..(114)
133 <223> OTHER INFORMATION: TTR
136 <220> FEATURE:
137 <221> NAME/KEY: variation

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138 <222> LOCATION: (145)..(147)
139 <223> OTHER INFORMATION: TTR
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144 <222> LOCATION: (151)..(153)
145 <223> OTHER INFORMATION: TTR
148 <220> FEATURE:
149 <221> NAME/KEY: variation
150 <222> LOCATION: (169)..(171)
151 <223> OTHER INFORMATION: AGR
154 <220> FEATURE:
155 <221> NAME/KEY: variation
156 <222> LOCATION: (172)..(174)
157 <223> OTHER INFORMATION: AGY
160 <220> FEATURE:
161 <221> NAME/KEY: variation
162 <222> LOCATION: (175)..(177)
163 <223> OTHER INFORMATION: AGR
166 <220> FEATURE:
167 <221> NAME/KEY: variation
168 <222> LOCATION: (244)..(246)
169 <223> OTHER INFORMATION: TTR
172 <220> FEATURE:
173 <221> NAME/KEY: variation
174 <222> LOCATION: (247)..(249)
175 <223> OTHER INFORMATION: TTR
178 <220> FEATURE:
179 <221> NAME/KEY: variation
180 <222> LOCATION: (265)..(267)
181 <223> OTHER INFORMATION: TTR
184 <220> FEATURE:
185 <221> NAME/KEY: variation
186 <222> LOCATION: (268)..(270)
187 <223> OTHER INFORMATION: AGY
190 <220> FEATURE:
191 <221> NAME/KEY: variation
192 <222> LOCATION: (289)..(291)
193 <223> OTHER INFORMATION: AGR
196 <220> FEATURE:
197 <221> NAME/KEY: variation
198 <222> LOCATION: (301)..(303)
199 <223> OTHER INFORMATION: TTR
202 <220> FEATURE:
203 <221> NAME/KEY: variation
204 <222> LOCATION: (310)..(312)
205 <223> OTHER INFORMATION: TTR
208 <220> FEATURE:
209 <221> NAME/KEY: variation
210 <222> LOCATION: (361)..(363)

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Input Set : A:\58764.000062.txt

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211 <223> OTHER INFORMATION: TTR
214 <220> FEATURE:
215 <221> NAME/KEY: variation
216 <222> LOCATION: (370)..(372)
217 <223> OTHER INFORMATION: TTR
220 <220> FEATURE:
221 <221> NAME/KEY: variation
222 <222> LOCATION: (409)..(411)
223 <223> OTHER INFORMATION: TTR
226 <220> FEATURE:
227 <221> NAME/KEY: variation
228 <222> LOCATION: (424)..(426)
229 <223> OTHER INFORMATION: TTR
232 <220> FEATURE:
233 <221> NAME/KEY: variation
234 <222> LOCATION: (436)..(438)
235 <223> OTHER INFORMATION: AGY
238 <220> FEATURE:
239 <221> NAME/KEY: variation
240 <222> LOCATION: (439)..(441)
241 <223> OTHER INFORMATION: TTR
244 <220> FEATURE:
245 <221> NAME/KEY: variation
246 <222> LOCATION: (490)..(492)
247 <223> OTHER INFORMATION: AGY
250 <220> FEATURE:
251 <221> NAME/KEY: variation
252 <222> LOCATION: (502)..(504)
253 <223> OTHER INFORMATION: AGY
256 <220> FEATURE:
257 <221> NAME/KEY: variation
258 <222> LOCATION: (511)..(513)
259 <223> OTHER INFORMATION: TTR
262 <220> FEATURE:
263 <221> NAME/KEY: variation
264 <222> LOCATION: (523)..(525)
265 <223> OTHER INFORMATION: AGY
268 <220> FEATURE:
269 <221> NAME/KEY: variation
270 <222> LOCATION: (550)..(552)
271 <223> OTHER INFORMATION: TTR
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275 <221> NAME/KEY: variation
276 <222> LOCATION: (562)..(564)
277 <223> OTHER INFORMATION: TTR
280 <220> FEATURE:
281 <221> NAME/KEY: variation
282 <222> LOCATION: (565)..(567)
283 <223> OTHER INFORMATION: AGR

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Input Set : A:\58764.000062.txt

Output Set: N:\CRF4\05312006\J580076.raw

286 <220> FEATURE:
287 <221> NAME/KEY: variation
288 <222> LOCATION: (580)..(582)
289 <223> OTHER INFORMATION: TTR
292 <220> FEATURE:
293 <221> NAME/KEY: variation
294 <222> LOCATION: (631)..(633)
295 <223> OTHER INFORMATION: AGY
298 <220> FEATURE:
299 <221> NAME/KEY: variation
300 <222> LOCATION: (637)..(639)
301 <223> OTHER INFORMATION: AGY
304 <220> FEATURE:
305 <221> NAME/KEY: variation
306 <222> LOCATION: (643)..(645)
307 <223> OTHER INFORMATION: TTR
310 <220> FEATURE:
311 <221> NAME/KEY: variation
312 <222> LOCATION: (658)..(660)
313 <223> OTHER INFORMATION: TTR
316 <220> FEATURE:
317 <221> NAME/KEY: variation
318 <222> LOCATION: (673)..(675)
319 <223> OTHER INFORMATION: TTR
322 <220> FEATURE:
323 <221> NAME/KEY: variation
324 <222> LOCATION: (697)..(699)
325 <223> OTHER INFORMATION: TTR
328 <220> FEATURE:
329 <221> NAME/KEY: variation
330 <222> LOCATION: (712)..(714)
331 <223> OTHER INFORMATION: AGY
334 <220> FEATURE:
335 <221> NAME/KEY: variation
336 <222> LOCATION: (715)..(717)
337 <223> OTHER INFORMATION: AGY
340 <220> FEATURE:
341 <221> NAME/KEY: variation
342 <222> LOCATION: (727)..(729)
343 <223> OTHER INFORMATION: TTR
346 <220> FEATURE:
347 <221> NAME/KEY: misc_feature
348 <222> LOCATION: (12)..(12)
349 <223> OTHER INFORMATION: N = A, G, C or T
352 <220> FEATURE:
353 <221> NAME/KEY: misc_feature
354 <222> LOCATION: (15)..(15)
355 <223> OTHER INFORMATION: N = A, G, C or T
358 <220> FEATURE:

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/580,076

DATE: 05/31/2006
TIME: 13:56:02

Input Set : A:\58764.000062.txt
Output Set: N:\CRF4\05312006\J580076.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; N Pos. 6, 11, 15, 27, 33, 54, 63, 66, 69, 75, 81, 84, 99, 105, 114, 135, 138, 144
Seq#:2; N Pos. 147, 153, 156, 162, 171, 174, 177, 184, 192, 225, 240, 246, 249, 264, 267
Seq#:2; N Pos. 270, 273, 276, 291, 294, 303, 306, 312, 315, 321, 327, 330, 336, 351, 363
Seq#:2; N Pos. 366, 372, 381, 396, 402, 411, 414, 426, 429, 432, 438, 441, 444, 465, 468
Seq#:2; N Pos. 492, 495, 504, 510, 513, 519, 525, 531, 543, 552, 555, 561, 564, 567, 582
Seq#:2; N Pos. 594, 615, 633, 638, 645, 660, 668, 675, 681, 699, 702, 708, 714, 717, 723
Seq#:2; N Pos. 729

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:2,3,4,5,6,7

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Input Set : A:\58764.000062.txt

Output Set: N:\CRF4\05312006\J580076.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:911 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:913 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:60
L:915 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:120
L:917 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:180
L:919 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:240
L:921 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:300
L:923 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:360
L:925 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:420
L:927 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:480
L:929 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:540
L:931 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:600
L:933 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:660
L:935 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:720